RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/573,885A

Source: IEW

Date Processed by STIC: 4/20/07

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IFWO

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        RAW SEQUENCE LISTING                    DATE: 04/20/2007
        PATENT APPLICATION:  US/10/573,885A     TIME: 16:01:24

        Input Set : E:\seqSU0411PCT.txt
        Output Set: N:\CRF4\04202007\J573885A.raw

    3 <110> APPLICANT: SUNTORY LIMITED
    5 <120> TITLE OF INVENTION: A gene encoding an enzyme catalyzing biosynthesis of lignan,
    6       and the use thereof
    8 <130> FILE REFERENCE: SU0411/PCT
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/573,885A
C--> 10 <141> CURRENT FILING DATE: 2006-03-29
   10 <150> PRIOR APPLICATION NUMBER: JP 2003-341313
   11 <151> PRIOR FILING DATE: 2003-09-30
   13 <150> PRIOR APPLICATION NUMBER: JP 2003-432383
   14 <151> PRIOR FILING DATE: 2003-12-26
   16 <160> NUMBER OF SEQ ID NOS: 79
   18 <170> SOFTWARE: PatentIn Ver. 2.1
   20 <210> SEQ ID NO: 1
   21 <211> LENGTH: 506
   22 <212> TYPE: PRT
   23 <213> ORGANISM: Sesamum indicum
   25 <220> FEATURE:
   26 <223> OTHER INFORMATION: SiP189
   28 <400> SEQUENCE: 1
   29 Met Glu Ala Glu Met Leu Tyr Ser Ala Leu Ala Leu Thr Phe Ala Ile
   30   1               5                  10                  15
   32 Phe Met Val Tyr Arg Ile Leu Ser Asn Ser Gln Asp Lys Arg Ser Leu
   33              20                  25                  30
   35 Thr Lys Leu Pro Pro Ser Pro Pro Gly Trp Leu Pro Val Ile Gly His
   36          35                  40                  45
   38 Ala His Leu Met Lys Asn Leu Leu His Arg Thr Leu Tyr Asp Phe Ser
   39      50                  55                  60
   41 Gln Lys Leu Gly Pro Ile Phe Ser Ile Arg Phe Gly Ser Arg Leu Val
   42 65                  70                  75                  80
   44 Val Val Val Ser Ser Ser Ser Leu Val Glu Glu Cys Phe Thr Lys Tyr
   45                  85                  90                  95
   47 Asp Ile Val Leu Ala Asn Arg Pro Gln Ala Ser Val Asp Arg Arg Ser
   48              100                 105                 110
   50 Leu Gly Phe Ser Thr Thr Ser Val Ile Gly Ala Pro Tyr Gly Asp His
   51          115                 120                 125
   53 Trp Arg Asn Leu Arg Lys Leu Cys Asp Leu Glu Val Phe Ala Pro Thr
   54          130                 135                 140
   56 Arg Leu Ala Ser Phe Leu Ser Ile Arg Leu Asp Glu Arg Asp Arg Met
   57 145                 150                 155                 160
   59 Ile Ser Ala Leu Tyr Lys Ile Ser Ser Ala Gly Phe Ala Lys Val Asn
   60              165                 170                 175
   62 Leu Glu Ala Lys Ile Val Glu Leu Thr Phe Asn Asn Ile Met Arg Met
   63              180                 185                 190
```

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65 Val Ala Ala Lys Arg Tyr Tyr Gly Glu Glu Ala Glu Asp Asp Glu Glu
66          195               200               205
68 Ala Lys Arg Phe Arg Asp Leu Thr Lys Glu Ala Leu Glu Leu Thr Ser
69     210               215               220
71 Ala Ser Asn Pro Gly Glu Ile Phe Pro Ile Leu Arg Trp Leu Gly Cys
72 225               230               235               240
74 Asn Gly Leu Glu Lys Lys Leu Ala Val His Ser Arg Lys Thr Asp Glu
75               245               250               255
77 Phe Met Gln Gly Leu Leu Asp Glu His Arg Arg Gly Glu Arg Gln Asn
78          260               265               270
80 Thr Met Val Asp His Leu Leu Ser Leu Gln Glu Ser Gln Pro Glu Tyr
81          275               280               285
83 Tyr Thr Asp Glu Ile Ile Thr Gly Leu Ile Val Ala Leu Ile Ile Ala
84     290               295               300
86 Gly Thr Asp Ala Ser Val Val Thr Thr Glu Trp Ala Met Ser Leu Leu
87 305               310               315               320
89 Leu Asn His Pro Lys Val Leu Glu Lys Ala Arg Lys Glu Leu Asp Thr
90               325               330               335
92 Leu Val Gly His Glu Arg Met Val Asp Glu His Asp Leu Pro Lys Leu
93          340               345               350
95 Arg Tyr Leu His Cys Ile Val Leu Glu Thr Leu Arg Leu Phe Pro Ser
96          355               360               365
98 Val Pro Thr Leu Val Pro His Glu Pro Ser Glu Asp Cys Lys Ile Gly
99     370               375               380
101 Gly Tyr Asn Val Pro Lys Gly Thr Met Val Leu Val Asn Ala Trp Ala
102 385               390               395               400
104 Ile His Arg Asp Pro Lys Val Trp Asp Asp Pro Leu Ser Phe Lys Pro
105               405               410               415
107 Asp Arg Phe Glu Ile Met Glu Val Glu Thr His Lys Leu Leu Pro Phe
108          420               425               430
110 Gly Met Gly Arg Arg Ala Cys Pro Gly Ala Gly Leu Ala Gln Lys Phe
111          435               440               445
113 Val Gly Leu Ala Leu Gly Ser Leu Ile Gln Cys Phe Asp Trp Glu Arg
114     450               455               460
116 Thr Ser Pro Glu Lys Ile Asp Leu Asn Glu Gly Ser Gly Ile Thr Leu
117 465               470               475               480
119 Pro Lys Ala Lys Thr Leu Glu Ala Met Cys Lys Pro Arg His Val Met
120               485               490               495
122 Glu Lys Val Leu Arg Gln Val Ser Asn Val
123          500               505
126 <210> SEQ ID NO: 2
127 <211> LENGTH: 1518
128 <212> TYPE: DNA
129 <213> ORGANISM: Sesamum indicum
131 <220> FEATURE:
132 <223> OTHER INFORMATION: SiP189
134 <400> SEQUENCE: 2
135 atggaagctg aaatgctata ttcagctctc gctctcacct tcgccatatt catggtttac 60
136 agaattcttt ctaattcgca ggacaagcgc agcctgacta agctgcctcc gagcccgccc 120
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Input Set : E:\seqSU0411PCT.txt
Output Set: N:\CRF4\04202007\J573885A.raw

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137 ggttggctgc cggtgatcgg ccacgctcat ctcatgaaaa atctcctcca tagaacacta 180
138 tacgacttct cccagaaact gggacccata ttttccatcc ggttcgggtc gcgcctcgtg 240
139 gtggtggtgt cctcctcctc cctggtggag gaatgtttca ccaagtatga cattgtcttg 300
140 gcaaatcgcc ctcaggcttc tgttgaccgg cgctcacttg ggttcagcac caccagcgta 360
141 atcggggccc cgtacgggga ccattggcgc aacctgcgaa agttgtcgga tcttgaagta 420
142 ttcgccccga cccgtctcgc ctcgttttta tccatcaggc ttgacgagag ggaccgcatg 480
143 atttccgcgt tatacaaaat ctcgtccgcc ggtttcgcga aggtgaattt ggaagcgaag 540
144 attgtggagc tgacgtttaa taacataatg aggatggtgg cggcgaagag atactatggg 600
145 gaggaggcgg aggacgacga ggaggcgaag aggttcaggg acctgacgaa ggaggctttg 660
146 gagttgacga gcgcttccaa tcctggtgag atatttccaa tattgcggtg gcttggttgc 720
147 aatgggctgg agaagaagct ggctgttcac tcgcggaaga cggatgagtt catgcaaggg 780
148 ctgctggacg aacaccgacg gggcgagcgc cagaacacca tggttgatca tttgctttcg 840
149 ttgcaggaat ctcaacctga gtactacact gatgaaatca tcactggcct catagttgca 900
150 ttgataattg cgggaacgga tgcatcggtt gtaactacag aatgggcgat gtcccttta 960
151 ctaaatcatc ccaaagtact tgaaaaggct agaaaagaac tggacactct agtaggacac 1020
152 gaacgcatgg ttgatgaaca cgatctcccc aaactacgtt accttcactg catagtcttg 1080
153 gagaccttaa ggttattccc ttctgttcca actttggtgc cacacgaacc atcagaggat 1140
154 tgtaaaattg ggggatacaa tgtccccaag gggacaatgg tattagtgaa tgcttgggca 1200
155 atacaccgag accccaaggt gtgggacgac cccttgagct ttaagcccga caggtttgag 1260
156 ataatggaag tggagacaca caagttgttg ccgttcggaa tgggcaggag agcgtgtcct 1320
157 ggagctggac tggcgcagaa gtttgtgggg ttggctttgg ggtcgctgat tcagtgtttc 1380
158 gactgggaga gaacgagtcc cgagaaaatt gacttgaacg aaggttctgg gataaccttg 1440
159 cctaaagcta agacgttgga agccatgtgc aaacctagac atgtcatgga aaaagttctt 1500
160 cgtcaggttt ccaacgtt                                              1518
163 <210> SEQ ID NO: 3
164 <211> LENGTH: 21
165 <212> TYPE: DNA
166 <213> ORGANISM: Artificial Sequence
168 <220> FEATURE:
169 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificially
170       Synthesized Primer Sequence, Si18SrRNA-FW
172 <400> SEQUENCE: 3
173 tatgcttgtc tcaaagatta a                                            21
176 <210> SEQ ID NO: 4
177 <211> LENGTH: 21
178 <212> TYPE: DNA
179 <213> ORGANISM: Artificial Sequence
181 <220> FEATURE:
182 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificially
183       Synthesized Primer Sequence, Si18SrRNA-RV
185 <400> SEQUENCE: 4
186 aacatctaag ggcatcacag a                                            21
189 <210> SEQ ID NO: 5
190 <211> LENGTH: 24
191 <212> TYPE: DNA
192 <213> ORGANISM: Artificial Sequence
194 <220> FEATURE:
195 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificially
196       Synthesized Primer Sequence, CYP90A-FW
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RAW SEQUENCE LISTING   DATE: 04/20/2007
PATENT APPLICATION: US/10/573,885A   TIME: 16:01:24

Input Set : **E:\seqSU0411PCT.txt**
Output Set: **N:\CRF4\04202007\J573885A.raw**

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198 <400> SEQUENCE: 5
199 ttttccgatg aagagattgt tgac                                    24
202 <210> SEQ ID NO: 6
203 <211> LENGTH: 18
204 <212> TYPE: DNA
205 <213> ORGANISM: Artificial Sequence
207 <220> FEATURE:
208 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificially
209      Synthesized Primer Sequence, CYP90A-RV
211 <400> SEQUENCE: 6
212 tgccatctcc aagggttg                                           18
215 <210> SEQ ID NO: 7
216 <211> LENGTH: 24
217 <212> TYPE: DNA
218 <213> ORGANISM: Artificial Sequence
220 <220> FEATURE:
221 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificially
222      Synthesized Primer Sequence, CYP72B-FW
224 <400> SEQUENCE: 7
225 cttaatgttc aaatgataat ggat                                    24
228 <210> SEQ ID NO: 8
229 <211> LENGTH: 18
230 <212> TYPE: DNA
231 <213> ORGANISM: Artificial Sequence
233 <220> FEATURE:
234 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificially
235      Synthesized Primer Sequence, CYP72B-RV
237 <400> SEQUENCE: 8
238 gtaaatcgtt cagggttg                                           18
241 <210> SEQ ID NO: 9
242 <211> LENGTH: 24
243 <212> TYPE: DNA
244 <213> ORGANISM: Artificial Sequence
246 <220> FEATURE:
247 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificially
248      Synthesized Primer Sequence, CYP71B-FW
250 <400> SEQUENCE: 9
251 ttcaccactg atcatctcaa agga                                    24
254 <210> SEQ ID NO: 10
255 <211> LENGTH: 18
256 <212> TYPE: DNA
257 <213> ORGANISM: Artificial Sequence
259 <220> FEATURE:
260 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificially
261      Synthesized Primer Sequence, CYP71B-RV
263 <400> SEQUENCE: 10
264 agaaacctgt cagggtta                                           18
267 <210> SEQ ID NO: 11
268 <211> LENGTH: 24
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269 <212> TYPE: DNA
270 <213> ORGANISM: Artificial Sequence
272 <220> FEATURE:
273 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificially
274       Synthesized Primer Sequence, CYP84A-FW
276 <400> SEQUENCE: 11
277 cttacccgtg acaatatcaa agca                                          24
280 <210> SEQ ID NO: 12
281 <211> LENGTH: 18
282 <212> TYPE: DNA
283 <213> ORGANISM: Artificial Sequence
285 <220> FEATURE:
286 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificially
287       Synthesized Primer Sequence, CYP84A-RV
289 <400> SEQUENCE: 12
290 aaaaacctcg atggtcta                                                 18
293 <210> SEQ ID NO: 13
294 <211> LENGTH: 24
295 <212> TYPE: DNA
296 <213> ORGANISM: Artificial Sequence
298 <220> FEATURE:
299 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificially
300       Synthesized Primer Sequence, CYP96A-FW
302 <400> SEQUENCE: 13
303 agtcatgata agttcctcag ggac                                          24
306 <210> SEQ ID NO: 14
307 <211> LENGTH: 18
308 <212> TYPE: DNA
309 <213> ORGANISM: Artificial Sequence
311 <220> FEATURE:
312 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificially
313       Synthesized Primer Sequence, CYP96A-RV
315 <400> SEQUENCE: 14
316 atccatctct ctggcttg                                                 18
319 <210> SEQ ID NO: 15
320 <211> LENGTH: 24
321 <212> TYPE: DNA
322 <213> ORGANISM: Artificial Sequence
324 <220> FEATURE:
325 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificially
326       Synthesized Primer Sequence, CYP710A-FW
328 <400> SEQUENCE: 15
329 tccgaagacg aagccatcgg cggt                                          24
332 <210> SEQ ID NO: 16
333 <211> LENGTH: 18
334 <212> TYPE: DNA
335 <213> ORGANISM: Artificial Sequence
337 <220> FEATURE:
338 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificially
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 04/20/2007
PATENT APPLICATION: US/10/573,885A   TIME: 16:01:25

Input Set : **E:\seqSU0411PCT.txt**
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<u>Please Note:</u>
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:77; N Pos. 2199

VERIFICATION SUMMARY   DATE: 04/20/2007
PATENT APPLICATION: US/10/573,885A   TIME: 16:01:25

            Input Set : E:\seqSU0411PCT.txt
            Output Set: N:\CRF4\04202007\J573885A.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1814 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77 after pos.:2160